

INFORMATION DISCLOSURE CITATION

Atty. Docket No.	7414.0025-01	Appln. No.	09/911,903
Applicant	Hugh J. Pasika et al.		
Filing Date	July 23, 2001	Group:	0661 1631

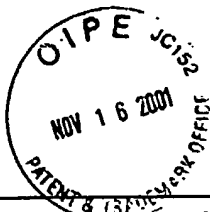
U.S. PATENT DOCUMENTS						
Examiner Initial*	Document Number	Issue Date	Name	Class	Subclass	Filing Date (if appropriate)
cm	5,580,728	Dec 3, 1996	Perlin			
1	6,019,896	Feb 1, 2000	Mansfield et al.			

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GROUP 1631
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FOREIGN PATENT DOCUMENTS						
	Document Number	Publication Date	Country	Class	Sub Class	Translation Yes or No
ca	WO 99/53423	Oct 21, 1999	PCT			

OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages, Etc.)	
ca	Ng, See-Kiong, "Automating Computational Molecular Genetics: Solving the Microsatellite Genotyping Problem,:" School of Computer Science, Carnegie Mellon University, Pittsburgh, PA 15213, Jan 23, 1998, submitted in partial fulfillment of the requirements for the degree of Doctor of Philosophy, pages 107-150.
	Pálsson, Birgir, et al., "Using Quality Measures to Facilitate Allele Calling in High-Throughput Genotyping," <u>Genome Research</u> , Vol. 9, No. 10, Oct 1999, pages 1002-1012.
	Perlin, Mark W., "An Expert System for Scoring DNA Database Profiles," Genetic Identity Conference Proceedings, 11 th International Symposium on Human Identification - 2000, Oct 7-14, 2000, Biloxi, MS, 30 pages.

Examiner	<i>altur</i>	Date Considered	4/16/04
*Examiner: Initial if reference considered, whether or not citation is in conformance with MPEP 609; draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.			
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OMB No. 0651-0011

Atty. Docket No.	7414-0025-01	Serial No.	09/911,903
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U.S. PATENT DOCUMENTS						
Examiner Initial*	Document Number	Issue Date	Name	Class	Sub Class	Filing Date If Appropriate
COL	5,541,067	Jul 30, 1996	Perlin			
	5,580,728	Dec 3, 1996	Perlin			
	5,853,979	Dec 29, 1998	Green et al.			
	5,871,628	Feb 16, 1999	Dabiri et al.			
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COL	ABI PRISM® Genotyper® 2.5 User's Manual, PE Biosystems, 1998, Foster City, CA.
COL	"Genetic Profiler Software," [online], [Retrieved on 5/15/2001]. Retrieved from http://www.mdyn.com/products/genotyping/software.htm , 3 pages.
	"Genotyping: High-throughput Genotyping," 5/15/2001, http://www.apbiotech.com/product/publication/lsn/lsn4/lsn4-1.html , 4 pages.
	Hoffman, E.P. et al., 1993, "Fully Automated PCR-based Disease Gene Mapping," <u>Am. J. Hum. Genet.</u> , 53(3 Supplement):Abstract 1172.
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	"MegaBACE Genotyping System Tools," 2000, <u>Amersham Pharmacia Biotech Inc.</u> , 2 pages.
	Ng, S.-K. et al., 1996, "Fully Automated Microsatellite Genotyping: Computer Software," <u>Am. J. Hum. Genet.</u> , 59(4 Supplement):Abstract 1319.
	"Savitzky-Golay Smoothing Filters", <u>Numerical Recipes in C: The Art of Scientific Computing</u> , 1992, Chapter 14.8, Press, William H. et al., Cambridge University Press, 650-655.
	Pálsson, B. et al., 1999, "Using Quality Measure to Facilitate Allele Calling in High-Throughput Genotyping," <u>Genome Research</u> , 9:1002-1012.
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cd	Perlin, M. W. et al., 1994, "Automated Genotyping of Dinucleotide Repeat Markers," <u>Am J. Hum. Genet.</u> , 55(3 Supplement):Abstract 1159.
	Perlin, M. W. et al., 1995, "Toward Fully Automated Genotyping: Genotyping Microsatellite Markers by Deconvolution," <u>Am. J. Hum. Genet.</u> , 57:1199-1210.
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